

TeX version 5.1.4_6_4578
copyright (c) 1992 TeX Corporation Ltd.

[illegible]

born on:
May 29, 2002,
at 22:16, 30 weeks
(with a term)
255,796 Milligram

15-19-125-548-16

[illegible]

61. http://www.bbc.co.uk/1/health/2004/04/040414_1_1.shtml

```
Series: table; IDENTITY=NO;
      (0.01, 0.01, 0.01)
      (0.01, 0.01, 0.01)
```

Submitted: 16/154066-5042, 8/10/77, 4/76, 4/75, 4/74, 4/73, 4/72, 4/71, 4/70, 4/69, 4/68, 4/67, 4/66, 4/65, 4/64, 4/63, 4/62, 4/61, 4/60, 4/59, 4/58, 4/57, 4/56, 4/55, 4/54, 4/53, 4/52, 4/51, 4/50, 4/49, 4/48, 4/47, 4/46, 4/45, 4/44, 4/43, 4/42, 4/41, 4/40, 4/39, 4/38, 4/37, 4/36, 4/35, 4/34, 4/33, 4/32, 4/31, 4/30, 4/29, 4/28, 4/27, 4/26, 4/25, 4/24, 4/23, 4/22, 4/21, 4/20, 4/19, 4/18, 4/17, 4/16, 4/15, 4/14, 4/13, 4/12, 4/11, 4/10, 4/9, 4/8, 4/7, 4/6, 4/5, 4/4, 4/3, 4/2, 4/1, 4/0, 3/99, 3/98, 3/97, 3/96, 3/95, 3/94, 3/93, 3/92, 3/91, 3/90, 3/89, 3/88, 3/87, 3/86, 3/85, 3/84, 3/83, 3/82, 3/81, 3/80, 3/79, 3/78, 3/77, 3/76, 3/75, 3/74, 3/73, 3/72, 3/71, 3/70, 3/69, 3/68, 3/67, 3/66, 3/65, 3/64, 3/63, 3/62, 3/61, 3/60, 3/59, 3/58, 3/57, 3/56, 3/55, 3/54, 3/53, 3/52, 3/51, 3/50, 3/49, 3/48, 3/47, 3/46, 3/45, 3/44, 3/43, 3/42, 3/41, 3/40, 3/39, 3/38, 3/37, 3/36, 3/35, 3/34, 3/33, 3/32, 3/31, 3/30, 3/29, 3/28, 3/27, 3/26, 3/25, 3/24, 3/23, 3/22, 3/21, 3/20, 3/19, 3/18, 3/17, 3/16, 3/15, 3/14, 3/13, 3/12, 3/11, 3/10, 3/9, 3/8, 3/7, 3/6, 3/5, 3/4, 3/3, 3/2, 3/1, 2/99, 2/98, 2/97, 2/96, 2/95, 2/94, 2/93, 2/92, 2/91, 2/90, 2/89, 2/88, 2/87, 2/86, 2/85, 2/84, 2/83, 2/82, 2/81, 2/80, 2/79, 2/78, 2/77, 2/76, 2/75, 2/74, 2/73, 2/72, 2/71, 2/70, 2/69, 2/68, 2/67, 2/66, 2/65, 2/64, 2/63, 2/62, 2/61, 2/60, 2/59, 2/58, 2/57, 2/56, 2/55, 2/54, 2/53, 2/52, 2/51, 2/50, 2/49, 2/48, 2/47, 2/46, 2/45, 2/44, 2/43, 2/42, 2/41, 2/40, 2/39, 2/38, 2/37, 2/36, 2/35, 2/34, 2/33, 2/32, 2/31, 2/30, 2/29, 2/28, 2/27, 2/26, 2/25, 2/24, 2/23, 2/22, 2/21, 2/20, 2/19, 2/18, 2/17, 2/16, 2/15, 2/14, 2/13, 2/12, 2/11, 2/10, 2/9, 2/8, 2/7, 2/6, 2/5, 2/4, 2/3, 2/2, 2/1, 1/99, 1/98, 1/97, 1/96, 1/95, 1/94, 1/93, 1/92, 1/91, 1/90, 1/89, 1/88, 1/87, 1/86, 1/85, 1/84, 1/83, 1/82, 1/81, 1/80, 1/79, 1/78, 1/77, 1/76, 1/75, 1/74, 1/73, 1/72, 1/71, 1/70, 1/69, 1/68, 1/67, 1/66, 1/65, 1/64, 1/63, 1/62, 1/61, 1/60, 1/59, 1/58, 1/57, 1/56, 1/55, 1/54, 1/53, 1/52, 1/51, 1/50, 1/49, 1/48, 1/47, 1/46, 1/45, 1/44, 1/43, 1/42, 1/41, 1/40, 1/39, 1/38, 1/37, 1/36, 1/35, 1/34, 1/33, 1/32, 1/31, 1/30, 1/29, 1/28, 1/27, 1/26, 1/25, 1/24, 1/23, 1/22, 1/21, 1/20, 1/19, 1/18, 1/17, 1/16, 1/15, 1/14, 1/13, 1/12, 1/11, 1/10, 1/9, 1/8, 1/7, 1/6, 1/5, 1/4, 1/3, 1/2, 1/1, 0/99, 0/98, 0/97, 0/96, 0/95, 0/94, 0/93, 0/92, 0/91, 0/90, 0/89, 0/88, 0/87, 0/86, 0/85, 0/84, 0/83, 0/82, 0/81, 0/80, 0/79, 0/78, 0/77, 0/76, 0/75, 0/74, 0/73, 0/72, 0/71, 0/70, 0/69, 0/68, 0/67, 0/66, 0/65, 0/64, 0/63, 0/62, 0/61, 0/60, 0/59, 0/58, 0/57, 0/56, 0/55, 0/54, 0/53, 0/52, 0/51, 0/50, 0/49, 0/48, 0/47, 0/46, 0/45, 0/44, 0/43, 0/42, 0/41, 0/40, 0/39, 0/38, 0/37, 0/36, 0/35, 0/34, 0/33, 0/32, 0/31, 0/30, 0/29, 0/28, 0/27, 0/26, 0/25, 0/24, 0/23, 0/22, 0/21, 0/20, 0/19, 0/18, 0/17, 0/16, 0/15, 0/14, 0/13, 0/12, 0/11, 0/10, 0/9, 0/8, 0/7, 0/6, 0/5, 0/4, 0/3, 0/2, 0/1, -1/99, -1/98, -1/97, -1/96, -1/95, -1/94, -1/93, -1/92, -1/91, -1/90, -1/89, -1/88, -1/87, -1/86, -1/85, -1/84, -1/83, -1/82, -1/81, -1/80, -1/79, -1/78, -1/77, -1/76, -1/75, -1/74, -1/73, -1/72, -1/71, -1/70, -1/69, -1/68, -1/67, -1/66, -1/65, -1/64, -1/63, -1/62, -1/61, -1/60, -1/59, -1/58, -1/57, -1/56, -1/55, -1/54, -1/53, -1/52, -1/51, -1/50, -1/49, -1/48, -1/47, -1/46, -1/45, -1/44, -1/43, -1/42, -1/41, -1/40, -1/39, -1/38, -1/37, -1/36, -1/35, -1/34, -1/33, -1/32, -1/31, -1/30, -1/29, -1/28, -1/27, -1/26, -1/25, -1/24, -1/23, -1/22, -1/21, -1/20, -1/19, -1/18, -1/17, -1/16, -1/15, -1/14, -1/13, -1/12, -1/11, -1/10, -1/9, -1/8, -1/7, -1/6, -1/5, -1/4, -1/3, -1/2, -1/1, -2/99, -2/98, -2/97, -2/96, -2/95, -2/94, -2/93, -2/92, -2/91, -2/90, -2/89, -2/88, -2/87, -2/86, -2/85, -2/84, -2/83, -2/82, -2/81, -2/80, -2/79, -2/78, -2/77, -2/76, -2/75, -2/74, -2/73, -2/72, -2/71, -2/70, -2/69, -2/68, -2/67, -2/66, -2/65, -2/64, -2/63, -2/62, -2/61, -2/60, -2/59, -2/58, -2/57, -2/56, -2/55, -2/54, -2/53, -2/52, -2/51, -2/50, -2/49, -2/48, -2/47, -2/46, -2/45, -2/44, -2/43, -2/42, -2/41, -2/40, -2/39, -2/38, -2/37, -2/36, -2/35, -2/34, -2/33, -2/32, -2/31, -2/30, -2/29, -2/28, -2/27, -2/26, -2/25, -2/24, -2/23, -2/22, -2/21, -2/20, -2/19, -2/18, -2/17, -2/16, -2/15, -2/14, -2/13, -2/12, -2/11, -2/10, -2/9, -2/8, -2/7, -2/6, -2/5, -2/4, -2/3, -2/2, -2/1, -3/99, -3/98, -3/97, -3/96, -3/95, -3/94, -3/93, -3/92, -3/91, -3/90, -3/89, -3/88, -3/87, -3/86, -3/85, -3/84, -3/83, -3/

Total number of hits satisfying chosen parameters: 3230142

[illegible]

Post-process	Min Match (%)	Max Match (%)	Listing	first 1000	summaries
--------------	---------------	---------------	---------	------------	-----------

1. *Abstracts* : *EST* : *

```
l: m_estb:*
```

2: en_estim:*

```
}; em_estim; *
```

4: em_estm: *

```
5: em_estov: *
6: em_estob: *
```

```

6: em_estro: *
7: em_estro: *

```

```

x: emmc:*
x: emmc:*
```

9: qb_011.*

```

10: qb_est2: *
11:
12:

```

[illegible]

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
---	---	---	---	---	---	---	---	---	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	-----

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
---	---	---	---	---	---	---	---	---	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	-----

[illegible]

1. Williamston *

17: qb_uss: *

ix: en-us-finn

ALLI-SCHMIDT : 61

```

21: emss_plm
22: emss_vlm
23: emss_vlm

```

14.4-551-116	1
14.4-551-116	2
14.4-551-116	3
14.4-551-116	4
14.4-551-116	5
14.4-551-116	6
14.4-551-116	7
14.4-551-116	8
14.4-551-116	9
14.4-551-116	10
14.4-551-116	11
14.4-551-116	12
14.4-551-116	13
14.4-551-116	14
14.4-551-116	15
14.4-551-116	16
14.4-551-116	17
14.4-551-116	18
14.4-551-116	19
14.4-551-116	20
14.4-551-116	21
14.4-551-116	22
14.4-551-116	23
14.4-551-116	24
14.4-551-116	25
14.4-551-116	26
14.4-551-116	27
14.4-551-116	28
14.4-551-116	29
14.4-551-116	30
14.4-551-116	31
14.4-551-116	32
14.4-551-116	33
14.4-551-116	34
14.4-551-116	35
14.4-551-116	36
14.4-551-116	37
14.4-551-116	38
14.4-551-116	39
14.4-551-116	40
14.4-551-116	41
14.4-551-116	42
14.4-551-116	43
14.4-551-116	44
14.4-551-116	45
14.4-551-116	46
14.4-551-116	47
14.4-551-116	48
14.4-551-116	49
14.4-551-116	50
14.4-551-116	51
14.4-551-116	52
14.4-551-116	53
14.4-551-116	54
14.4-551-116	55
14.4-551-116	56
14.4-551-116	57
14.4-551-116	58
14.4-551-116	59
14.4-551-116	60
14.4-551-116	61
14.4-551-116	62
14.4-551-116	63
14.4-551-116	64
14.4-551-116	65
14.4-551-116	66
14.4-551-116	67
14.4-551-116	68
14.4-551-116	69
14.4-551-116	70
14.4-551-116	71
14.4-551-116	72
14.4-551-116	73
14.4-551-116	74
14.4-551-116	75
14.4-551-116	76
14.4-551-116	77
14.4-551-116	78
14.4-551-116	79
14.4-551-116	80
14.4-551-116	81
14.4-551-116	82
14.4-551-116	83
14.4-551-116	84
14.4-551-116	85
14.4-551-116	86
14.4-551-116	87
14.4-551-116	88
14.4-551-116	89
14.4-551-116	90
14.4-551-116	91
14.4-551-116	92
14.4-551-116	93
14.4-551-116	94
14.4-551-116	95
14.4-551-116	96
14.4-551-116	97
14.4-551-116	98
14.4-551-116	99
14.4-551-116	100

[illegible]
$$\text{std_dev} = 1.7$$

25; em_qss_othe

26: em_ssh_pro

27: em_qss_red:

THE

Plot. No. is the number of
store greater than or equal

and is derived by analysis

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARY

[illegible]

16.1	86.3	804	12	B641741B	HVSMEK001	15.8	83.2	456	17	AV646209	AV646209
16.2	86.3	804	13	B6517684	G6282580	15.8	83.2	456	17	AV646209	AV646209
16.3	86.3	804	14	B67008967	MTL00190	15.8	83.2	456	17	AV646209	AV646209
16.4	86.3	804	15	B6442766	01 R L80	15.8	83.2	456	17	AV646209	AV646209
16.5	86.3	804	16	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
16.6	86.3	804	17	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
16.7	86.3	804	18	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
16.8	86.3	804	19	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
16.9	86.3	804	20	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.0	86.3	804	21	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.1	86.3	804	22	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.2	86.3	804	23	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.3	86.3	804	24	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.4	86.3	804	25	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.5	86.3	804	26	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.6	86.3	804	27	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.7	86.3	804	28	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.8	86.3	804	29	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
17.9	86.3	804	30	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.0	86.3	804	31	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.1	86.3	804	32	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.2	86.3	804	33	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.3	86.3	804	34	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.4	86.3	804	35	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.5	86.3	804	36	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.6	86.3	804	37	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.7	86.3	804	38	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.8	86.3	804	39	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
18.9	86.3	804	40	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
19.0	86.3	804	41	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
19.1	86.3	804	42	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
19.2	86.3	804	43	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
19.3	86.3	804	44	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209
19.4	86.3	804	45	B6524056	602925745	15.8	83.2	456	17	AV646209	AV646209

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89</											

```

538 15.4 81.1 534 13 R1901904 fd15k01.y
539 15.4 81.1 535 17 AZ218729 Sheared D
540 15.4 81.1 541 17 R1901647 fd15k01.x
541 15.4 81.1 542 17 AA014148 mb191111.r
542 15.4 81.1 543 17 R0575257 K D330036
543 15.4 81.1 544 17 AQ421105 K0C1-11-1
544 15.4 81.1 545 17 R05292584 603102158
545 15.4 81.1 546 17 RH224127 1006116H1
546 15.4 81.1 547 17 AQ644771 KPC193-Ec
547 15.4 81.1 548 17 R0556220 U1-M-RH0-
548 15.4 81.1 549 17 CR6501 CR6501.H008
549 15.4 81.1 550 17 A1125382 ms510517.r
550 15.4 81.1 551 17 R1584117 602057717
551 15.4 81.1 552 17 AA197831 ms04405.r
552 15.4 81.1 553 17 R0738116 K PST0992
553 15.4 81.1 554 17 R0510446 R0201016
554 15.4 81.1 555 17 R0701917 U1-E-Q01-
555 15.4 81.1 556 17 RE333056 us43d11.y
556 15.4 81.1 557 17 R1159850 602863650
557 15.4 81.1 558 17 R0476378 602521884
558 15.4 81.1 559 17 R1196699 R1196699
559 15.4 81.1 560 17 R0080200 R3050H06-
560 15.4 81.1 561 17 AZ782980 280024116
561 15.4 81.1 562 17 R0291787 601084554
562 15.4 81.1 563 17 RE996452 U1-M-C00p
563 15.4 81.1 564 17 A0139397 AU139397
564 15.4 81.1 565 17 R0510927 TH4-GH030
565 15.4 81.1 566 17 R0462904 U1-M-C00p
566 15.4 81.1 567 17 R0372729 6030071
567 15.4 81.1 568 17 R0334475 us79002.y
568 15.4 81.1 569 17 R1987819 7212-21.H
569 15.4 81.1 570 17 R0897403 HGAL3.1.F
570 15.4 81.1 571 17 R1043233 R3043233
571 15.4 81.1 572 17 R1660064 R1660064
572 15.4 81.1 573 17 AV391710 AV391710
573 15.4 81.1 574 17 R0717861 U1-P-E09-
574 15.4 81.1 575 17 A1071572 w1a1061.x
575 15.4 81.1 576 17 R1648776 604277308
576 15.4 81.1 577 17 R0578535 602092978
577 15.4 81.1 578 17 R0367259 R3052D01-
578 15.4 81.1 579 17 R03974152 604943969
579 15.4 81.1 580 17 R120664 0254087.y
580 15.4 81.1 581 17 R0504859 10474007.y
581 15.4 81.1 582 17 R0774550 K-P33006.r
582 15.4 81.1 583 17 R0335735 S81-0035
583 15.4 81.1 584 17 R1036335 603037117
584 15.4 81.1 585 17 R0734913 601224337
585 15.4 81.1 586 17 R0766280 MY-13-0-0
586 15.4 81.1 587 17 R1555543 604339352
587 15.4 81.1 588 17 R0694009 1000152.H
588 15.4 81.1 589 17 R0909302 KPC1.24.3
589 15.4 81.1 590 17 R1705098 603070082
590 15.4 81.1 591 17 R1917737 603183827
591 15.4 81.1 592 17 R1722102 603059185
592 15.4 81.1 593 17 R145617 602910646
593 15.4 81.1 594 17 R0591313 AV594212
594 15.4 81.1 595 17 R1661967 603040459
595 15.4 81.1 596 17 R081108 601271186
596 15.4 81.1 597 17 R162495 601768832
597 15.4 81.1 598 17 A0139779 AU139779
598 15.4 81.1 599 17 R1103981 602890270
599 15.4 81.1 600 17 R053535 60381145
600 15.4 81.1 601 17 R0283122 60276133
601 15.4 81.1 602 17 R0285687 602380729
602 15.4 81.1 603 17 R1091848 602958718
603 15.4 81.1 604 17 R1693275 603343578
604 15.4 81.1 605 17 R1914315 603180975
605 15.4 81.1 606 17 R1767242 602957934
606 15.4 81.1 607 17 R1754873 604026853
607 15.4 81.1 608 17 R0514244 604033529
608 15.4 81.1 609 17 R1821091 604033551
609 15.4 81.1 610 17 AL543599 AL543599
534 15.4 81.1 764 13 R1901904 fd15k01.y
535 15.4 81.1 765 13 AZ218729 Sheared D
536 15.4 81.1 766 13 R1901647 fd15k01.x
537 15.4 81.1 767 13 AA014148 mb191111.r
538 15.4 81.1 768 13 R0575257 K D330036
539 15.4 81.1 769 13 AQ421105 K0C1-11-1
540 15.4 81.1 770 13 R05292584 603102158
541 15.4 81.1 771 13 RH224127 1006116H1
542 15.4 81.1 772 13 AQ644771 KPC193-Ec
543 15.4 81.1 773 13 R0556220 U1-M-RH0-
544 15.4 81.1 774 13 CR6501 CR6501.H008
545 15.4 81.1 775 13 A1125382 ms510517.r
546 15.4 81.1 776 13 R1584117 602057717
547 15.4 81.1 777 13 AA197831 ms04405.r
548 15.4 81.1 778 13 R0738116 K PST0992
549 15.4 81.1 779 13 R0510446 R0201016
550 15.4 81.1 780 13 R0701917 U1-E-Q01-
551 15.4 81.1 781 13 RE333056 us43d11.y
552 15.4 81.1 782 13 R1159850 602863650
553 15.4 81.1 783 13 R0476378 602521884
554 15.4 81.1 784 13 R1196699 R1196699
555 15.4 81.1 785 13 R0080200 R3050H06-
556 15.4 81.1 786 13 AZ782980 280024116
557 15.4 81.1 787 13 R0291787 601084554
558 15.4 81.1 788 13 RE996452 U1-M-C00p
559 15.4 81.1 789 13 A0139397 AU139397
560 15.4 81.1 790 13 R0510927 TH4-GH030
561 15.4 81.1 791 13 R0462904 U1-M-C00p
562 15.4 81.1 792 13 R0372729 6030071
563 15.4 81.1 793 13 R0334475 us79002.y
564 15.4 81.1 794 13 R1987819 7212-21.H
565 15.4 81.1 795 13 R0897403 HGAL3.1.F
566 15.4 81.1 796 13 R1043233 R3043233
567 15.4 81.1 797 13 R1660064 R1660064
568 15.4 81.1 798 13 AV391710 AV391710
569 15.4 81.1 799 13 R0717861 U1-P-E09-
570 15.4 81.1 800 13 A1071572 w1a1061.x
571 15.4 81.1 801 13 R1648776 604277308
572 15.4 81.1 802 13 R0578535 602092978
573 15.4 81.1 803 13 R0367259 R3052D01-
574 15.4 81.1 804 13 R03974152 604943969
575 15.4 81.1 805 13 R120664 0254087.y
576 15.4 81.1 806 13 R0504859 10474007.y
577 15.4 81.1 807 13 R0774550 K-P33006.r
578 15.4 81.1 808 13 R0335735 S81-0035
579 15.4 81.1 809 13 R1036335 603037117
580 15.4 81.1 810 13 R0734913 601224337
581 15.4 81.1 811 13 R0766280 MY-13-0-0
582 15.4 81.1 812 13 R1555543 604339352
583 15.4 81.1 813 13 R0694009 1000152.H
584 15.4 81.1 814 13 R0909302 KPC1.24.3
585 15.4 81.1 815 13 R1705098 603070082
586 15.4 81.1 816 13 R1917737 603183827
587 15.4 81.1 817 13 R1722102 603059185
588 15.4 81.1 818 13 R145617 602910646
589 15.4 81.1 819 13 R0591313 AV594212
590 15.4 81.1 820 13 R1661967 603040459
591 15.4 81.1 821 13 R081108 601271186
592 15.4 81.1 822 13 R162495 601768832
593 15.4 81.1 823 13 A0139779 AU139779
594 15.4 81.1 824 13 R1103981 602890270
595 15.4 81.1 825 13 R053535 60381145
596 15.4 81.1 826 13 R0283122 60276133
597 15.4 81.1 827 13 R0285687 602380729
598 15.4 81.1 828 13 R1091848 602958718
599 15.4 81.1 829 13 R1693275 603343578
600 15.4 81.1 830 13 R1914315 603180975
601 15.4 81.1 831 13 R1767242 602957934
602 15.4 81.1 832 13 R1754873 604026853
603 15.4 81.1 833 13 R0514244 604033529
604 15.4 81.1 834 13 R1821091 604033551
605 15.4 81.1 835 13 AL543599 AL543599

```

15.3	81.1	94.9	14	160976649	6255514	ADEN500R1	7.87	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	94.2	12	161167961	6255514	ADEN500R1	7.89	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	95.3	12	161766964	6255514	ADEN500R1	7.91	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	95.3	13	162511888	6255514	ADEN500R1	7.91	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	95.5	13	162929289	6255514	ADEN500R1	7.91	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	95.9	12	163084296	6255514	ADEN500R1	7.92	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	96.1	12	163641643	6255514	ADEN500R1	7.93	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	96.2	17	163804088	6255514	ADEN500R1	7.93	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	96.9	14	164649450	6255514	ADEN500R1	7.95	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	96.9	14	164669773	6255514	ADEN500R1	7.95	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	96.9	14	164947714	6255514	ADEN500R1	7.95	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	97.2	12	165334360	6255514	ADEN500R1	7.96	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	98.2	14	165798150	6255514	ADEN500R1	7.97	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	99.2	12	165798352	6255514	ADEN500R1	7.97	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	99.8	14	166058444	6255514	ADEN500R1	7.97	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	99.4	13	166233985	6255514	ADEN500R1	7.97	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	100.0	14	166983687	6255514	ADEN500R1	7.98	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	100.4	14	166983687	6255514	ADEN500R1	7.98	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	101.0	14	166983687	6255514	ADEN500R1	7.98	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	101.0	14	166983687	6255514	ADEN500R1	7.98	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	101.0	17	168020118	6255514	ADEN500R1	7.98	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	101.7	14	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	ADEN500R1	7.99	15	72.9	6.71	17	7295367	ADEN500R1
15.3	81.1	102.3	13	168779706	6255514	AD							

810	14.8	77.9	414	13	B005251	B005251	B005251	887	11.8	77.9	587	13	AM25055	AM25055
811	14.8	77.9	421	14	B057572	B057572	B057572	888	14.8	77.9	588	14	B057572	B057572
812	14.8	77.9	422	15	B071433	B071433	B071433	889	14.8	77.9	589	15	B071433	B071433
813	14.8	77.9	425	9	A010121	A010121	A010121	890	14.8	77.9	590	9	A010121	A010121
814	14.8	77.9	425	9	A010121	A010121	A010121	891	14.8	77.9	591	9	A010121	A010121
815	14.8	77.9	425	10	AW95186	AW95186	AW95186	892	14.8	77.9	592	10	AW95186	AW95186
816	14.8	77.9	426	9	A010121	A010121	A010121	893	14.8	77.9	593	9	A010121	A010121
817	14.8	77.9	428	9	A010121	A010121	A010121	894	14.8	77.9	594	9	A010121	A010121
818	14.8	77.9	428	10	AW95186	AW95186	AW95186	895	14.8	77.9	595	10	AW95186	AW95186
819	14.8	77.9	430	14	B00762	B00762	B00762	896	14.8	77.9	596	14	B00762	B00762
820	14.8	77.9	431	14	B057572	B057572	B057572	897	14.8	77.9	597	14	B057572	B057572
821	14.8	77.9	432	17	A010121	A010121	A010121	898	14.8	77.9	598	17	A010121	A010121
822	14.8	77.9	432	17	A010121	A010121	A010121	899	14.8	77.9	599	17	A010121	A010121
823	14.8	77.9	434	10	B003163	B003163	B003163	900	14.8	77.9	600	10	B003163	B003163
824	14.8	77.9	435	9	A010121	A010121	A010121	901	14.8	77.9	601	9	A010121	A010121
825	14.8	77.9	437	13	B057572	B057572	B057572	902	14.8	77.9	602	13	B057572	B057572
826	14.8	77.9	439	13	B057572	B057572	B057572	903	14.8	77.9	603	13	B057572	B057572
827	14.8	77.9	440	14	B034014	B034014	B034014	904	14.8	77.9	604	14	B034014	B034014
828	14.8	77.9	444	12	B051624	B051624	B051624	905	14.8	77.9	605	12	B051624	B051624
829	14.8	77.9	445	17	A010121	A010121	A010121	906	14.8	77.9	606	17	A010121	A010121
830	14.8	77.9	446	17	A010121	A010121	A010121	907	14.8	77.9	607	17	A010121	A010121
831	14.8	77.9	449	5	A010121	A010121	A010121	908	14.8	77.9	608	5	A010121	A010121
832	14.8	77.9	450	9	A010121	A010121	A010121	909	14.8	77.9	609	9	A010121	A010121
833	14.8	77.9	452	12	B071433	B071433	B071433	910	14.8	77.9	610	12	B071433	B071433
834	14.8	77.9	454	14	B031036	B031036	B031036	911	14.8	77.9	611	14	B031036	B031036
835	14.8	77.9	455	10	B026438	B026438	B026438	912	14.8	77.9	612	10	B026438	B026438
836	14.8	77.9	456	9	A010121	A010121	A010121	913	14.8	77.9	613	9	A010121	A010121
837	14.8	77.9	459	12	B028138	B028138	B028138	914	14.8	77.9	614	12	B028138	B028138
838	14.8	77.9	459	17	A029694	A029694	A029694	915	14.8	77.9	615	17	A029694	A029694
839	14.8	77.9	463	14	B000483	B000483	B000483	916	14.8	77.9	616	14	B000483	B000483
840	14.8	77.9	464	17	A035156	A035156	A035156	917	14.8	77.9	617	17	A035156	A035156
841	14.8	77.9	465	14	B011874	B011874	B011874	918	14.8	77.9	618	14	B011874	B011874
842	14.8	77.9	466	12	B037648	B037648	B037648	919	14.8	77.9	619	12	B037648	B037648
843	14.8	77.9	471	10	B031179	B031179	B031179	920	14.8	77.9	620	10	B031179	B031179
844	14.8	77.9	471	14	B028361	B028361	B028361	921	14.8	77.9	621	14	B028361	B028361
845	14.8	77.9	473	10	B073977	B073977	B073977	922	14.8	77.9	622	10	B073977	B073977
846	14.8	77.9	475	9	A010121	A010121	A010121	923	14.8	77.9	623	9	A010121	A010121
847	14.8	77.9	477	13	B019782	B019782	B019782	924	14.8	77.9	624	13	B019782	B019782
848	14.8	77.9	479	9	A010121	A010121	A010121	925	14.8	77.9	625	9	A010121	A010121
849	14.8	77.9	479	14	B010521	B010521	B010521	926	14.8	77.9	626	14	B010521	B010521
850	14.8	77.9	482	17	A029792	A029792	A029792	927	14.8	77.9	627	17	A029792	A029792
851	14.8	77.9	483	10	A010121	A010121	A010121	928	14.8	77.9	628	10	A010121	A010121
852	14.8	77.9	483	10	A010121	A010121	A010121	929	14.8	77.9	629	10	A010121	A010121
853	14.8	77.9	486	10	A066504	A066504	A066504	930	14.8	77.9	630	10	A066504	A066504
854	14.8	77.9	491	13	B027871	B027871	B027871	931	14.8	77.9	631	13	B027871	B027871
855	14.8	77.9	491	13	B049852	B049852	B049852	932	14.8	77.9	632	13	B049852	B049852
856	14.8	77.9	492	14	B026029	B026029	B026029	933	14.8	77.9	633	14	B026029	B026029
857	14.8	77.9	493	9	A010121	A010121	A010121	934	14.8	77.9	634	9	A010121	A010121
858	14.8	77.9	493	13	B027871	B027871	B027871	935	14.8	77.9	635	13	B027871	B027871
859	14.8	77.9	494	14	B04384	B04384	B04384	936	14.8	77.9	636	14	B04384	B04384
860	14.8	77.9	496	17	A010121	A010121	A010121	937	14.8	77.9	637	17	A010121	A010121
861	14.8	77.9	496	17	A010121	A010121	A010121	938	14.8	77.9	638	17	A010121	A010121
862	14.8	77.9	503	10	A010121	A010121	A010121	939	14.8	77.9	639	10	A010121	A010121
863	14.8	77.9	503	17	B058585	B058585	B058585	940	14.8	77.9	640	17	B058585	B058585
864	14.8	77.9	504	12	B044771	B044771	B044771	941	14.8	77.9	641	12	B044771	B044771
865	14.8	77.9	510	14	B057763	B057763	B057763	942	14.8	77.9	642	14	B057763	B057763
866	14.8	77.9	515	17	A029762	A029762	A029762	943	14.8	77.9	643	17	A029762	A029762
867	14.8	77.9	515	9	A010121	A010121	A010121	944	14.8	77.9	644	9	A010121	A010121
868	14.8	77.9	519	17	A010121	A010121	A010121	945	14.8	77.9	645	17	A010121	A010121
869	14.8	77.9	520	14	B044970	B044970	B044970	946	14.8	77.9	646	14	B044970	B044970
870	14.8	77.9	521	10	A010121	A010121	A010121	947	14.8	77.9	647	10	A010121	A010121
871	14.8	77.9	522	10	A010121	A010121	A010121	948	14.8	77.9	648	10	A010121	A010121
872	14.8	77.9	522	12	B054126	B054126	B054126	949	14.8	77.9	649	12	B054126	B054126
873	14.8	77.9	522	12	B054126	B054126	B054126	950	14.8	77.9	650	12	B054126	B054126
874	14.8	77.9	523	12	B073617	B073617	B073617	951	14.8	77.9	651	12	B073617	B073617
875	14.8	77.9	524	13	B021260	B021260	B021260	952	14.8	77.9	652	13	B021260	B021260
876	14.8	77.9	525	10	A010121	A010121	A010121	953	14.8	77.9	653	10	A010121	A010121
877	14.8	77.9	525	10	A010121	A010121	A010121	954	14.8	77.9	654	10	A010121	A010121
878	14.8	77.9	526	12	B048147	B048147	B048147	955	14.8	77.9	655	12	B048147	B048147
879	14.8	77.9	526	12	B048147	B048147	B048147	956	14.8	77.9	656	12	B048147	B048147
880	14.8	77.9	527	9	A010121	A010121	A010121	957	14.8	77.9	657	9	A010121	A010121
881	14.8	77.9	527	9	A010121	A010121	A010121	958	14.8	77.9	658	9	A010121	A010121
882	14.8	77.9	528	14	B020762	B020762	B020762	959	14.8	77.9	659	14	B020762	B020762

[illegible]

cDNA library preparation, life technologies, inc.
 cDNA library arrayed by: The I.M.A.G.E. Consortium (LMNI)
 DNA Sequencing by: Inverte Genomics, Inc.
 Clone distribution: MGC clone distribution information can be
 found through the I.M.A.G.E. Consortium/LMNI at:
<http://image.llnl.gov>

Plate: LLAM9740 row: a column: 07
 High quality sequence stop: 524.

FEATURES

Source:

1, 734
 Location/Qualifiers
 Zorandism "Homo sapiens"
 ZR_seq "Zorandism"
 Zclone "IMAGE:5912078"
 Zclone_lib "NHLMC271"
 Zissue_type "Telomerase RNA"
 Zlab_host "PHOB" (phase-resistant)
 Zvec "Zorandism" (phase-resistant)
 SITE_2: Salt: Cloned unidirectionally. Primer: column d1.
 Average insert size 2.1 kb.
 194 a 194 c 214 g 111 t 1 others

Query Match: 100.0% Score: 197 19 125 Length: 734.
 Best Local Similarity: 100.0% Ident. No.: 42002
 Matches: 19, Conserved: 0, Mismatches: 0, Indels: 0, Gaps: 0.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

RESULT 39
 BASE COUNT 176 a 204 c 224 g 154 t
 ORIGIN
 constructed by R. Strausberg, Ph.D.,
 Institute of Molecular Biology, National Institutes of Health, Bethesda, Maryland, U.S.A. This is a NIH library.

QY 1 CCWAGACAGAGGAGGCGC 19
 11111111111111111111

14 183 CCWAGACAGAGGAGGCGC 165

DEFINITION

Query Match 100.0%; Score 19; Db 13; Length 792;
Best local Similarity 100.0%; Pred. No. 4, 3, 0, 0, 2;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

27 1 CAAACAGACAGAGCGGCGTC 19
|||||
28 62 CAAACAGACAGCGGCGTC 64

RESULT 44

EST 1

DEFINITION

U760349 Homo sapiens mRNA linear EST 25-SEP-2001
603442291; REFSeq_116 Homo sapiens cDNA clone IMAGE7165567 5';
mRNA sequence.

ACCESSION

U760349

VERSION

U760349.1

KEYWORDS

EST.

SOURCE

ORGANISM

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Homidae; Homo;

1 (bases 1 to 796)

244 EST; EST; GenBank; RefSeq; Mammalia; Euteleostomi (M);

National Institutes of Health; National Institutes of Health (NIH);

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: eapbs@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA library Preparation: Life Technologies, Inc.

cDNA library Arrayed by: The L.M.A.G.E. Consortium (LME)

Cloned by: Invertebrate Genomics, Inc.

Clone Distribution: M62 clone distribution information can be

found through the L.M.A.G.E. Consortium/NIH at:

http://lmea1143.rockefeller.edu

Plate: LME1143, row 4, column 08

High quality sequence stop: 703.

Location/contig: 1, 796

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

DEFINITION

U760349

VERSION

U760349.1

KEYWORDS

EST.

SOURCE

ORGANISM

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Homidae; Homo;

1 (bases 1 to 815)

244 EST; EST; GenBank; RefSeq; Mammalia; Euteleostomi (M);

National Institutes of Health; National Institutes of Health (NIH);

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: eapbs@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA library Preparation: Life Technologies, Inc.

cDNA library Arrayed by: The L.M.A.G.E. Consortium (LME)

Cloned by: Invertebrate Genomics, Inc.

Clone Distribution: M62 clone distribution information can be

found through the L.M.A.G.E. Consortium/NIH at:

http://lmea1143.rockefeller.edu

Plate: LME1143, row 4, column 08

High quality sequence stop: 703.

Location/contig: 1, 815

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

DEFINITION

U760349

VERSION

U760349.1

KEYWORDS

EST.

SOURCE

ORGANISM

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Homidae; Homo;

1 (bases 1 to 815)

244 EST; EST; GenBank; RefSeq; Mammalia; Euteleostomi (M);

National Institutes of Health; National Institutes of Health (NIH);

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: eapbs@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA library Preparation: Life Technologies, Inc.

cDNA library Arrayed by: The L.M.A.G.E. Consortium (LME)

Cloned by: Invertebrate Genomics, Inc.

Clone Distribution: M62 clone distribution information can be

found through the L.M.A.G.E. Consortium/NIH at:

http://lmea1143.rockefeller.edu

Plate: LME1143, row 4, column 08

High quality sequence stop: 703.

Location/contig: 1, 815

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

Accession: "Homo sapiens"

